

SEQUENCE LISTING

10031327.022102

<110> Huang, Tim Hui-Ming

<120> HIGH-THROUGHPUT METHODS FOR DETECTING DNA METHYLATION

<130> 40629-2

<140> US 10/081,327

<141> 2002-02-21

<150> 09/497,855

<151> 2000-02-04

<150> 60/120,592

<151> 1999-02-18

<150> 60/118,760

<151> 1999-02-05

<160> 54

<170> PatentIn version 3.1

<210> 1

<211> 630

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(630)

<223> All n's are unknown

<400> 1

```
atgatttncg ctcttgata ttatcatctt gaacgaaagc tggtaaactc tggcttatgg      60
ataaaacaaa actttaccna tegtatttgg gtgagggggt ggggaatgga taactgtttc     120
ctctgctaaa attttagcgt ntnaatcggg ggatngagat ttgcataagt ttgacggggag     180
ggtcntgtnt tegtttcaag tatgggtnat tgtccacacc ttttcttnaa gantngcgcg      240
gggggcctaa ttcgaanggt naanaaaana aggaagacng angtgccana atgnacccca      300
atccacanct tatggganan ctcccaacgc gttngatgca aacttgaatn tcnntactnt      360
ccctaaatn ncttgngtt atentggeen aactgtttcc tgnnttaaaa tnnttancgg      420
ctccccattc cnetctannt acnaancggg aagcttnaat tttnaanncc nggggtnccct      480
aatganttaa ctaacccctt tanttncett nnncccnge cegnttcnat cgggaaactn      540
tttccccct ncnnttatta atcncncnc cccggaaaan gcggttgctt tntgggccct      600
nccctectcc caatganccc cncceggntt                                     630
```

<210> 2

<211> 913

<212> DNA